TAATCAAACTATGGAGAAGAAATCACTAGCTGGCTTATGCTTCCTCTTCTTGGTTCTCTTGTT M E K K S L A G L C F L F L V L F V

GCACAAGAAATTATGGTGACAGAAGCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGAC E Q E I M V T E A R T C E N L A D K Y R G P Δ

CATGCTTTAGTGGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGTGTAGCCFF S G C D T H C T T K E N A V S G R C R

GGACGACTTCCGCTGCTGGTGTACTAAAAGATGTTAAATGGATCTCCTCCAACATCAAGATGTG D D F R C W C T K R C *

CAAATAGTCTTTATAATAAAACTAAATAAATAAAATGCACGCAGTATAGCTACAACTTCATCTA

TTATATGTACTCAATATCGNGCATAACGTATTAGTTATGCACTTCTATCATATGGAATAAACAT

MEKKSLACLSFLLLVLFVAQETVVSEANTCENLAGSYKGVCFGGCDRHCRTQEGA I SGRCRDDFRCWCTKNC

| Alfafp2 Alfafp2 TGCTTCCTCTTCCTCGTTCTCTTGTTGAACAAGAAATTATGGTGACCGAG Alfafp1 TGCTTCCTCTTCTTGGTTCTCTTTGTTGAACAAGAAATTATGGTGACCGAG Alfafp1 TGCTTCCTCTTCTTGGTTCTCTTTGTTGCACAAGAAATTATGGTGACCGAG ******************************* | | AlfAFP2 ALFAFP1 |
|---|---|--------------------|
| Alfafp1 TGCTTCCTCTTCTTGGTTCTCTTTGTTGCACAAGAAATTGTGGTGACAGAA *********** * ********* * ********* * **** | 2 TATATTCATCAATCTAATCAAACTATGGAGAAGAAATCACTAGCTGGCT | |
| Alfafpl GCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGACCATGCTTTAGT ** ** ********* * * *** ********** ** Alfafpl GGTTGTGACTTTCACTGCAAAACCAAAGAACACTTACTTA | 1 TGCTTCCTCTTGGTTCTCTTTGTTGCACAAGAAATTGTGGTGACAG | |
| Alfafp1 GGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGTGT ********* ******* ******* * * * * * * | GCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGACCATGCTTTA | |
| Alfafpl AGGGACGACTTCCGCTGCTGCTGGATCC | GGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGT | |
| | AGGGACGACTTCCGCTGCTGGATCC | |

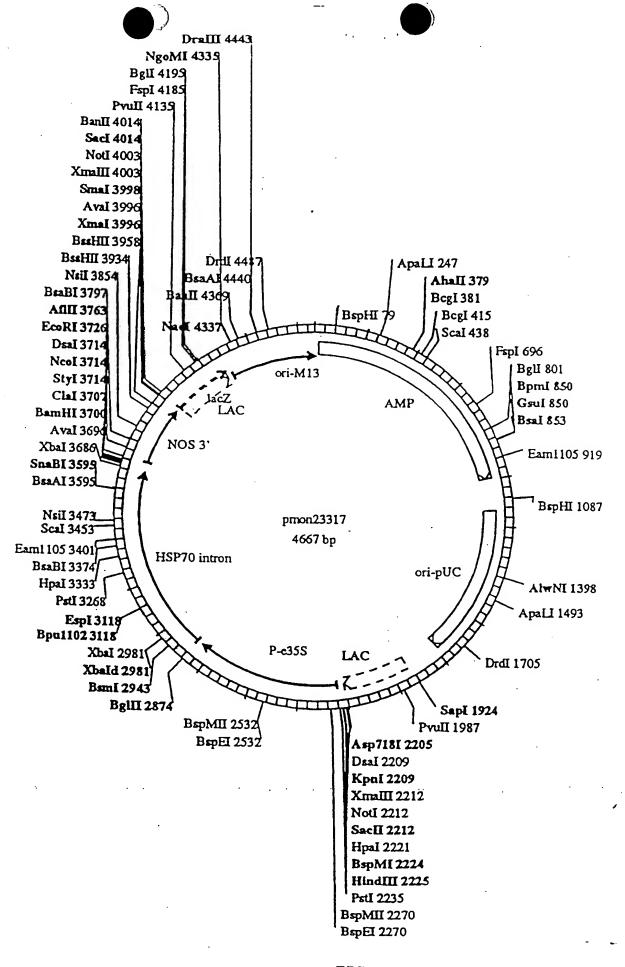


FIG. 4

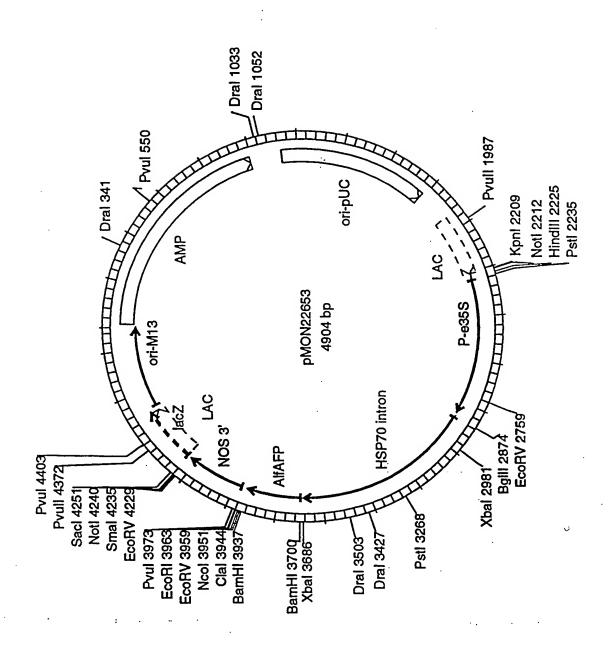


FIG. 5

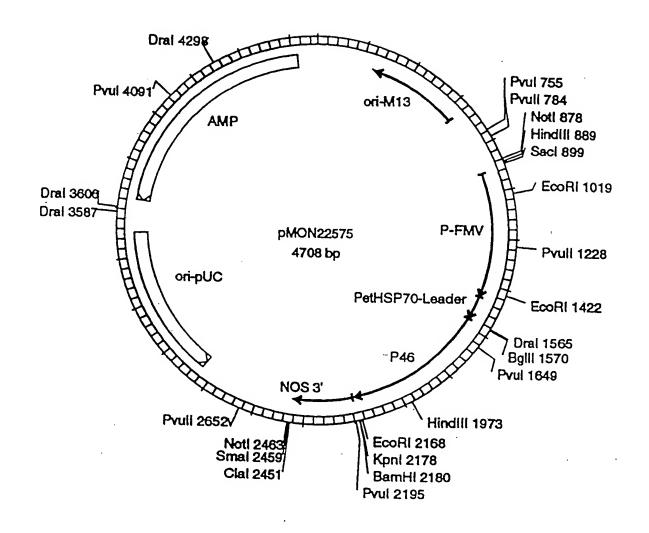


FIG. 6

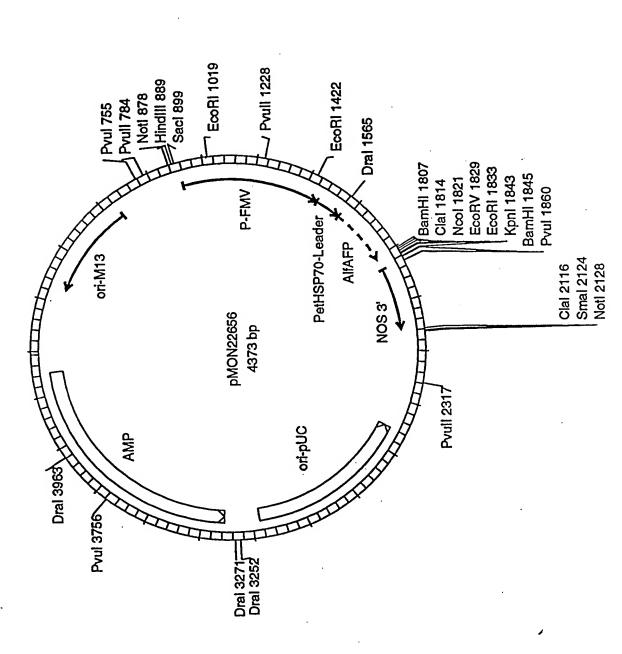


FIG. 7

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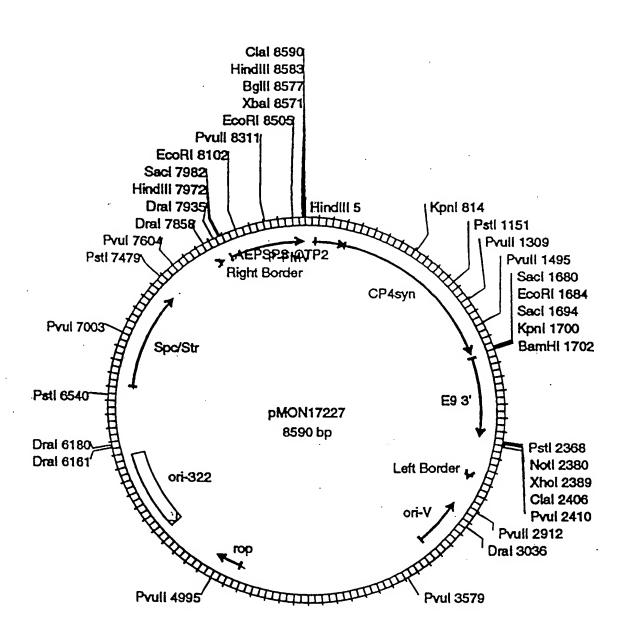


FIG. 8

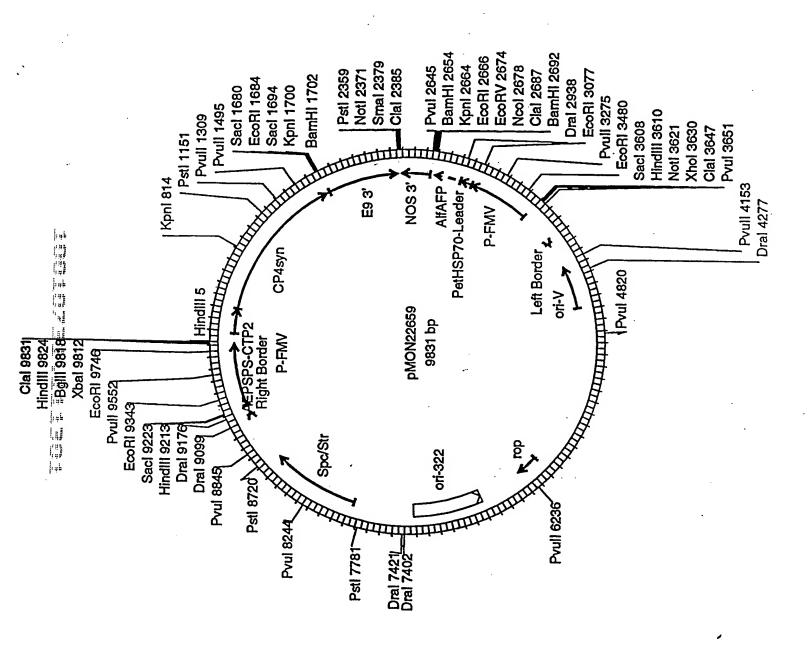
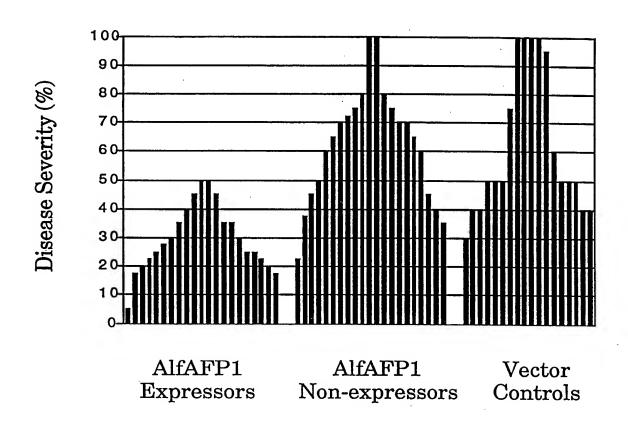


FIG. 9





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